

Query Match		100.0%	Score 1108;	DB 2;	Length 205;
Best Local Similarity		100.0%;	Pred. No. 2.2e-95;		
Matches 205;		Conservative	;	Mismatches	0;
				Indels	0;
				Gaps	0;
Qy	1	MWLLPALLLLCLSGCLSLKPGFSVITGAGSLTWQCYESMYGKYNKYWCRGQYDTSCE	60		
Db	1	MWLLPALLLLCLSGCLSLKPGFSVITGAGSLTWQCYESMYGKYNKYWCRGQYDTSCE	60		
Qy	61	IVETKGEKVERNGRVSIRDHPEALFTVTMQNLNEDDAGSYWCKIQYVWLVDWSRDF	120		
Db	61	IVETKGEKVERNGRVSIRDHPEALFTVTMQNLNEDDAGSYWCKIQYVWLVDWSRDF	120		
Qy	121	DLVRVYVSPAIIETPRRTTHATPPIELVNVNGRNLSIREVTIONSGFRLLSPHLLVLL	180		
Db	121	DLVRVYVSPAIIETPRRTTHATPPIELVNVNGRNLSIREVTIONSGFRLLSPHLLVLL	180		

Qy	181	KLPLLLSMLGAVFWVNRPQWAPPGR	205
Db	181	KLPLLLSMLGAVFWVNRPQWAPPGR	205

<!--EndFragment-->